## SEQUENCE LISTING

<110> KYOWA HAKKO KOGYO CO., LTD.

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<130> 11620W01.

<150> JP2003-350166

<151> 2003-10-09

<160> 32

 $\langle 170 \rangle$  Patent In Ver. 2.1

<210> 1

<211> 2008

<212> DNA

<213> Cricetulus griseus

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<213 Mus musculus

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<212> DNA

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<sup>⟨210⟩ 5</sup> 

<sup>&</sup>lt;211> 575

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Cricetulus griseus

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170

175

165

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185

190

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- His Val Glu Glu His Phe Gln Leu Leu Ala Arg Arg Met Gln Val Asp 385 390 395 400
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<213 > Homo sapiens

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Ser Ile Met Thr Asp Leu Tyr Tyr Leu Ser Gln Thr Asp Gly Ala Gly
35 40 45

Asp Trp Arg Glu Lys Glu Ala Lys Asp Leu Thr Glu Leu Val Gln Arg 50 55 60

Arg Ile Thr Tyr Leu Gln Asn Pro Lys Asp Cys Ser Lys Ala Lys Lys 165 70 75 80

Leu Val Cys Asn Ile Asn Lys Gly Cys Gly Tyr Gly Cys Gln Leu His
85 90 95

His Val Val Tyr Cys Phe Met Ile Ala Tyr Gly Thr Gln Arg Thr Leu 100 105 110

Ile	Leu	Glu	Ser	Gln	Asn	Trp	Arg	Tyr	Ala	Thr	Gly	Gly	Trp	Glu	Thr
		115					120					125.			

Val Phe Arg Pro Val Ser Glu Thr Cys Thr Asp Arg Ser Gly Ile Ser 130 135 140

Thr Gly His Trp Ser Gly Glu Val Lys Asp Lys Asn Val Gln Val Val 145 150 155 160

Glu Leu Pro Ile Val Asp Ser Leu His Pro Arg Pro Pro Tyr Leu Pro 165 170 175

Leu Ala Val Pro Glu Asp Leu Ala Asp Arg Leu Val Arg Val His Gly 180 185 190

Asp Pro Ala Val Trp Trp Val Ser Gln Phe Val Lys Tyr Leu Ile Arg 195 200 205

Pro Gln Pro Trp Leu Glu Lys Glu Ile Glu Glu Ala Thr Lys Lys Leu 210 215 220

Gly Phe Lys His Pro Val Ile Gly Val His Val Arg Arg Thr Asp Lys 225 230 235 240

Val Gly Thr Glu Ala Ala Phe His Pro Ile Glu Glu Tyr Met Val His 245 250 255

Val Glu Glu His Phe Gln Leu Leu Ala Arg Arg Met Gln Val Asp Lys 260 265 270

Lys Arg Val Tyr Leu Ala Thr Asp Asp Pro Ser Leu Leu Lys Glu Ala 275 280 285

Lys Thr Lys Tyr Pro Asn Tyr Glu Phe Ile Ser Asp Asn Ser Ile Ser 290 295 300 Trp Ser Ala Gly Leu His Asn Arg Tyr Thr Glu Asn Ser Leu Arg Gly 305 310 315 320

Val Ile Leu Asp Ile His Phe Leu Ser Gln Ala Asp Phe Leu Val Cys 325 330 335

Thr Phe Ser Ser Gln Val Cys Arg Val Ala Tyr Glu Ile Met Gln Thr 340 345 350

Leu His Pro Asp Ala Ser Ala Asn Phe His Ser Leu Asp Asp Ile Tyr 355 360 365

Tyr Phe Gly Gly Gln Asn Ala His Asn Gln Ile Ala Ile Tyr Ala His 370 375 380

Gln Pro Arg Thr Ala Asp Glu Ile Pro Met Glu Pro Gly Asp Ile Ile 385 390 395 400

Gly Val Ala Gly Asn His Trp Asp Gly Tyr Ser Lys Gly Val Asn Arg 405 410 415

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<211> 575

<212> PRT

<213> Sus scrofa

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Lys	Leu 50	Glu	Arg	Leu	Lys	Gln 55	Gln	Asn	Glu	Asp	Leu 60	Arg	Arg	Met	Alà
Glu 165	Ser	Leu	Arg	Ile	Pro 70	Glu	Gly	Pro	Ile	Asp 75	Gln	Gly	Pro	Ala	Ser 80
Gly	Arg	Val	Arg	Ala 85	Leu	Glu	Glu	Gln	Phe 90	Met	Lys	Ala	Lys	Glu 95	Gln
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Glu	Ile	Leu 115	Arg	Arg	Arg	Ile	Glu 120	Asn	Gly	Ala	Lys	Glu 125	Leu	Trp	Phe
Phe	Leu 130	Gln	Ser	Glu	Leu	Lys 135	Lys	Leu	Lys	Asn	Leu 140	Glu	Gly	Asn	Glu
Leu 145	Gln	Arg	His	Ala	Asp 150	Glu	Phe	Leu	Ser	Asp 155	Leu	Gly	His	His	Glu 160
Arg	Ser	Ile.	Met	Thr 165	Asp	Leu	Tyr		Leu 170	Ser	Gln	Thr	Asp	Gly 175	Ala
Gly	Asp	Trp	Arg 180	Glu	Lys	Glu	Ala	Lys 185	Asp	Leu	Thr	Glu	Leu 190	Val	Gln

Arg Arg Ile Thr Tyr Leu Gln Asn Pro Lys Asp Cys Ser Lys Ala Lys

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His 225	His	Val	Val	Tyr	Cys 230	Phe	Met	Ile	Ala	Tyr 235	Gly	Thr	Gln	Arg	Thr 240
Leu	Ala	Leu	Glu	Ser 245	His	Asn	Trp	Ar-g	Tyr 250	Ala	Thr	Gly	Gly	Trp 255	Glu
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Val	G1u 290	Leu	Pro	Ile	Val	Asp 295	Ser	Val	His	Pro	Arg 300	Pro	Pro	Tyr	Leu
Pro 305	Leu	Ala	Val	Pro	Glu 310	Asp	Leu	Ala	Asp	Arg 315	Leu	Val	Arg	Val	His 320
Gly	Asp	Pro	Ala	Val 325	Trp	Trp	Val	Ser	G1n 330	Phe	Val	Lys	Tyr	Leu 335	Ile
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Leu	Gly	Phe 355	Lys	His	Pro	Val	Ile 360	Gly	Val	His	Val	Arg 365	Arg	Thr	Asp
Lys	Val 370	Gly	Ala	Glu	Ala	Ala 375	Phe	His	Pro-	Ile	Glu 380	Glu	Tyr	Thr	Val

His Val Glu Glu Asp Phe Gln Leu Leu Ala Arg Arg Met Gln Val Asp

385					390					395					400
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Ser	Trp	Ser 435	Ala	Gly	Leu	His	Asn 440	Arg	Tyr	Thr	Glu	Asn 445	Ser	Leu	Arg
Gly	Val 450	Ile	Leu	Asp	Ile	His 455	Phe	Leu	Ser	Gln	Ala 460	Asp	Phe	Leu	Val
Cys 465	Thr	Phe	Ser	Ser	Gln 470	Val	Cys	Arg	Val	Ala 475	Tyr	Glu	Ile	Met	Gln 480
Ala	Leu	His	Pro	Asp 485	Ala	Ser	Ala	Asn	Phe 490	Arg	Ser	Leu	Asp	Asp 495	I.l e
Tyr	Tyr	Phe	Gly 500	Gly	Pro	Asn	Ala	His 505	Asn	Gln	Ile	Ala	Ile 510	Tyr	Pro
His	Gln	Pro-	Arg	Thr	Glu	Gly	Glu 520	Ile	Pro	Met	Glu	Pro 525	Gly	Asp	Ile
Ile	Gly 530	Val	Ala	Gly	Asn	His 535	Trp	Asp	Gly		Pro 540	Lys	Gly	Val	Asn
Arg 545		Leu	Gly	Arg	Thr 550	Gly	Leu	Tyr	Pro	Ser 555	Tyr	Lys	Val	Arg	Glu 560
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<211> 9196

<212> DNA

<213> Cricetulus griseus

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Val Ile Met Ser Arg Gly Gln Ile Val Leu Ser Gln Ser Pro Ala I	
20 25 30	
ctg tct gca tct cca ggg gag aag gtc aca atg act tgc agg gcc ag	gc 144
Leu Ser Ala Ser Pro Gly Glu Lys Val Thr Met Thr Cys Arg Ala Se	-
35 40 45	
tca agt gta agt tac atc cac tgg ttc cag cag aag cca gga tcc to	cc 192
Ser Ser Val Ser Tyr Ile His Trp Phe Gln Gln Lys Pro Gly Ser Se	
50 55 60	•
ccc aaa ccc tgg att tat gcc aca tcc aac ctg gct tct gga gtc cc	ct 240
Pro Lys Pro Trp Ile Tyr Ala Thr Ser Asn Leu Ala Ser Gly Val Pr	
	80

gtt cgc ttc agt ggc agt ggg tct ggg act t	tet tae tet ete ace ate 288
Val Arg Phe Ser Gly Ser Gly Ser Gly Thr S	Ser Tyr Ser Leu Thr Ile
85 90	. 95
agc aga gtg gag gct gaa gat gct gcc act t	at tac tgc cag cag tgg 336
Ser Arg Val Glu Ala Glu Asp Ala Ala Thr T	yr Tyr Cys Gln Gln Trp
100 105	110
act agt aac cca ccc acg ttc gga ggg ggg a	acc aag ctg gaa atc aaa 384
Thr Ser Asn Pro Pro Thr Phe Gly Gly Gly T	·
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1 5 10	15
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Val Leu Ser Gln Val Gln Leu Gln Gln Pro G	ly Ala Glu Leu Val Lys
20 25	30
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Pro Gly Ala Ser Val Lys Met Ser Cys Lys A	lla Ser Gly Tyr Thr Phe
35 40	45
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Thr Ser Tyr Asn Met His Trp Val Lys Gln T	hr Pro Gly Arg Gly Leu
50 55	60
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Glu Trp Ile Gly Ala Ile Tyr Pro Gly Asn G	ly Asp Thr Ser Tyr Asn
65 70	75 80
.cag aag ttc aaa ggc aag gcc aca ttg act g	gca gac aaa tcc tcc agc 288
Gln Lys Phe Lys Gly Lys Ala Thr Leu Thr A	· ·
85 90	95
aca gcc tac atg cag ctc agc agc ctg aca t	
and goo tan and had not ago ago ong and t	or gag gav for gog gre out

Thr	Ala	Tyr	Met	Gln	Leu	Ser			Thr	Ser	Glu	Asp		Ala	Val	
			100	•				105					110			
tat	tac	tgt	gca.	aga	tcg	ac t	tac	tac	ggc	ggt	gac	tgg	tac	ttc	aat	384
Tyr	Tyr	Cys.	Ala	Arg	Ser	Thr	Tyr	Tyr	Gly	Gly	Asp	Trp	Tyr	Phe	Asn	
	-	115					120					125				
gtc	tgg	ggc	gca	ggg	acc	acg	gtc	acc	gtc	tct	gca					420
Val	Trp	Gly	Ala	Gly	Thr	Thr	Val	Thr	Val	Ser	Ala					
	130					135	1.0				140			•		
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ctte	ctg	cta :	atcag	gtgc	tt ca	agtca	ataa	t g								91
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